

Please amend the application as follows:

In the claims:

The listing of claims below will replace all prior versions, and listings, of claims in the application.

1-31. (cancelled)

32. (currently amended) A method for screening compounds for modulation of GABA_A receptor 1 transcription, comprising the steps of:

(a) providing a host cell ~~hosting that has been transfected~~ with an expression system comprising a nucleic acid molecule constituting:

a promoter element selected from the group consisting of:

(i) a nucleic acid molecule comprising SEQ ID NO: 1,

(ii) a nucleic acid molecule at least 95% homologous to SEQ ID NO: 1,

(iii) a nucleic acid molecule comprising SEQ ID NO: 2, and

(iv) a nucleic acid molecule at least 95% homologous to SEQ ID NO: 2; and

a reporter gene, wherein the promoter element is coupled to the reporter gene so that expression of the reporter gene is under the control of the promoter element;

- (b) contacting a test compound with the cell; and
- (c) determining whether the test compound modulates the level of expression of the reporter gene.

33. (previously presented) The method according to claim 32, wherein the reporter gene is selected from the group consisting of:

- (a) the firefly luciferase gene;
- (b) the bacterial chloramphenicol acetyl transferase (CAT) gene;
- (c) the β -galactosidase (β -Gal) gene; and
- (d) the green fluorescent protein (GFP) gene.

34. (previously presented) The method according to claim 32, wherein the host cell endogenously expresses at least one GABA_A receptor 1.

35. (currently amended) The method according to claim 32,
wherein the host cell ~~hosts~~ has further been transfected with an
expression system comprising a nucleic acid molecule encoding at
least one specific transcription factor.

36. (currently amended) The method according to claim 35,
wherein the specific transcription factor is selected from the
group consisting of: CREB-1, CREB-2, CREM-1, ATF-1, ATF-2, ATF-
3, ATF-4, Sp1, Sp2, Sp3, Sp4, AP-1 and AP-2.

37. (currently amended) A method for screening compounds for
modulation of GABA_B receptor 1 transcription, comprising the
steps of:

(a) providing a host cell ~~hosting~~ that has been transfected
with an expression system comprising a nucleic acid molecule
constituting:

a promoter element consisting essentially of (1) a
functionally equivalent modified ~~form~~ variant of or (2) an
active fragment of a nucleic acid molecule selected from the
group consisting of:

(i) the nucleic acid molecule defined as SEQ ID NO: 1,
and

(ii) the nucleic acid molecule defined as SEQ ID NO:
2, and wherein the functionally equivalent modified form variant
of (i) above is at least 95% homologous to SEQ ID NO: 1 or SEQ
ID NO: 2; and

a reporter gene, wherein the promoter element is coupled to
the reporter gene so that expression of the reporter gene is
under the control of the promoter element;

(b) contacting a test compound with the cell; and

(c) determining whether the test compound modulates the
level of expression of the reporter gene.

38. (previously presented) The method according to claim 37,
wherein the reporter gene is selected from the group consisting
of:

(a) the firefly luciferase gene;

(b) the bacterial chloramphenicol acetyl transferase (CAT)
gene;

(c) the β -galactosidase (β -Gal) gene; and

(d) the green fluorescent protein (GFP) gene.

39. (previously presented) The method according to claim 37, wherein the host cell endogenously expresses at least one GABA_B receptor 1.

40. (currently amended) The method according to claim 37, wherein the host cell ~~hosts~~ has further been transfected with an expression system comprising a nucleic acid molecule encoding at least one specific transcription factor.

41. (currently amended) The method according to claim 40, wherein the specific transcription factor is selected from the group consisting of: CREB-1, CREB-2, CREM-1, ATF-1, ATF-2, ATF-3, ATF-4, Sp1, Sp2, Sp3, Sp4, AP-1 and AP-2.

42. (currently amended) A method for screening compounds for modulation of GABA_B receptor 1 transcription, comprising the steps of:

(a) providing a host cell hosting that has been transfected with an expression system comprising a nucleic acid molecule constituting:

a promoter element consisting essentially of (1) a functionally equivalent modified form variant of or (2) an active fragment of the nucleic acid molecule defined as SEQ ID NO: 1, the promoter element comprising:

(i) the nucleic acid sequence of positions 3009-3016 of SEQ ID NO: 1,

(ii) the nucleic acid sequence of positions 3037-3044 of SEQ ID NO: 1, and

(iii) the nucleic acid sequence of positions 3116-3123 of SEQ ID NO: 1,

and wherein the functionally equivalent modified form variant of (1) above is at least 95% homologous to SEQ ID NO: 1; and

a reporter gene, wherein the promoter element is coupled to the reporter gene so that expression of the reporter gene is under the control of the promoter element;

(b) contacting a test compound with the cell; and

(c) determining whether the test compound modulates the level of expression of the reporter gene.

43. (previously presented) The method according to claim 42, wherein the promoter element is not operably linked to a repressor region of a GABA_B receptor 1 Pla promoter.

44. (currently amended) A method for screening compounds for modulation of GABA_B receptor 1 transcription, comprising the steps of:

(a) providing a host cell hosting that has been transfected with an expression system comprising a nucleic acid molecule constituting:

a promoter element consisting essentially of (1) a functionally equivalent modified ~~form~~ variant of or (2) an active fragment of the nucleic acid molecule defined as SEQ ID NO: 2, the promoter element comprising the nucleic acid sequence of positions 4308-4315 of SEQ ID NO: 2

and wherein the functionally equivalent modified ~~form~~ variant of (1) above is at least 95% homologous to SEQ ID NO: 2, and

a reporter gene, wherein the promoter element is coupled to the reporter gene so that expression of the reporter gene is under the control of the promoter element;

(b) contacting a test compound with the cell; and

(c) determining whether the test compound modulates the level of expression of the reporter gene.

45. (previously presented) The method according to claim 44, wherein the promoter element further comprises:

(i) the nucleic acid sequence of positions 4080-4087 of SEQ ID NO: 2;

(ii) the nucleic acid sequence of positions 4196-4204 of SEQ ID NO: 2;

(iii) the nucleic acid sequence of positions 4241-4249 of SEQ ID NO: 2; and

(iv) the nucleic acid sequence of positions 4272-4279 of SEQ ID NO: 2.

46. (previously presented) The method according to claim 44, wherein the promoter element is not operably linked to a repressor region of a GABA_B receptor 1 P1b promoter.

47. (previously presented) The method according to claim 46,
wherein the promoter element further comprises:

(i) the nucleic acid sequence of positions 4080-4087 of SEQ
ID NO: 2;

(ii) the nucleic acid sequence of positions 4196-4204 of
SEQ ID NO: 2;

(iii) the nucleic acid sequence of positions 4241-4249 of
SEQ ID NO: 2; and

(iv) the nucleic acid sequence of positions 4272-4279 of
SEQ ID NO: 2.

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